Electricity Generation by a Phototrophic Bacterium in a Glucose−Fed Double Chambered Microbial Fuel Cell Using a Fabricated 3D Anode Electrode

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INTRODUCTION

Energy is essential for life in the twenty-first century, considering its significance in various sectors, as evidenced by the constant rise in global energy consumption. Since their discovery, fossil fuels have been the major global energy provider (more than 80%) (Alimonti, 2018). The intensifying efforts to extract more fossil fuels and their extensive applications due to increasing demand resulted in more carbon dioxide production, raising global concerns (Guo et al., 2022). According to the United Nations (UN), 75% of greenhouse gas (GHG) emissions result from fossil fuel use, consequently destabilizing the world’s environmental equilibrium and causing the global threat of climate change accelerated by continuous deforestation (Sonawane et al., 2017).
A significant increase in global energy demand has been experienced for the past decades. With the progressive increase in global energy demands, as shown in Figure 1, and the Sustainable Development Goals (SDGs) call for sustainable, affordable, and clean energy, the race for improving and innovating various techniques intensifies exponentially. Several renewable, sustainable, and eco-friendly energy alternatives like solar, wind, and hydropower have been developed and improved. Although these renewable energy technologies (RE) are now key contributors to the modern energy mix globally with little or no emissions of GHGs, their cost of installation/operation is noticeably expensive, making it unaffordable and inaccessible to nearly 775 million people around the world (International Renewable Energy Agency, IRENA, 2024; International Energy Agency, IEA, 2022).

Bioenergy results from utilizing organic, agricultural, and industrial wastes to generate energy with no or minimal GHG emissions (Roder and Welfle, 2019). Ghangrekar et al. (2022) stated bioelectricity is the most focused area in bioenergy. Bioelectricity involves converting the stored chemical energy in organic substances into electrical energy by microorganisms (aerobically or anaerobically).

In 1911, Potter introduced the idea of the Microbial Fuel Cell (MFC), an electrochemical cell with electrodes (anode and cathode) and chamber(s) containing the microorganism(s) and electrolytes (organic waste) to be degraded for bioelectricity generation (Potter 1991; Haruna et al., 2023). Potter’s initiative used *E. coli* and *Saccharomyces* with platinum electrodes to generate energy. Unfortunately, due to the insignificant amount of energy generated, researchers showed no further interest until 1931, when Cohen used a similar method to generate a potential voltage of 35 V (Logan, 2005). Over time, interest in this research area catapulted, with an average publication record of 1,244 articles annually between 2010−2022, according to the Web of Science (2023) database.

In addition to generating bioelectricity, MFCs provide an avenue for wastewater treatment, thereby reducing the volume of global wastewater (380 billion m$^3$), which is expected to increase by 51% by 2050, according to the European Investment Bank (EIB, 2022). Moreover, 80% of global infectious diseases are caused by untreated wastewater, resulting in 13 million deaths annually (Wolf et al., 2023).

The performance of an MFC is majorly dictated by the MFC configuration, microbial electrochemical activity, substrate type, electrode type, and electrode material (Pant et al., 2010; Kim et al., 2011). Several configurations such as mediator/mediatorless MFC, photoMFC, upflow MFC, and stacked−MFC have been made to improve MFCs’ performance, and that resulted in a significant increase in power density (PD) and current density (CD).
generation (Jang et al., 2004; Aelteman et al., 2006). In a recent study, Sato et al. (2023) incorporated an MFC with a hydroponic system (MFC–Hyp) to generate a maximum PD and CD of 523.9±50 mW/m² and 2,676.7±340.7 mA/m², respectively. Subsequently, another study used a photo-MFC configuration to generate a maximum PD of 280 mW/m² with photocathode and bioanode (Wang et al., 2023).

Furthermore, other developments were recorded by Shirkoshi et al. (2022) with microfluidic-MFC, which produced a maximum open circuit voltage (OCV) of 1.295 V, PD of 21,111 W/m³, and CD of 140,000 A/m³. Apollon et al. (2023a) also recorded a PD and CD of 46.97 ± 0.67 mW/m² and 77.45 ± 0.24 mA/m² using plant–based MFC clay embedded with Stevia rebandiana. While with sediment-MFC configuration, a significant PD of 1,056.6 W/m² was obtained by Mejia-Lopez et al. (2023). However, most configurations are on the research level, while some have been applied on an industrial scale. In this study, double-chambered MFC was used because of its availability, ease to scale-up, and efficiency (Wang et al., 2023).

Not only did electrode materials such as carbon, graphene, polyaniline, platinum, and stainless-steel fibre play a vital role in enhancing MFC performance, but also the type, i.e., three-dimensional (3D) and two-dimensional (2D) as well, since it influences microbial colonisation and hence the electrochemical activity of the microbe (Qi et al., 2018). Several studies have indicated electrode materials’ role in microbial adhesion and performance. Tan et al. (2018), using in-situ Co₃O₄ nanoparticles modified nitrogen–doped graphene (3D GN–Co₃O₄) cathode, recorded 578±10 mW/m² and 0.45±0.01 V as PD and OCV, respectively, which is fivefold higher compared to the 177.03 mW/m² generated using plain carbon cathode by Piskin and Genc (2023).

The differences in concentration and type of substrate also showed an improved energy generation in MFCs, as Tian et al. (2017) evaluated. Likewise, Guo et al. (2022) stated that an increase in electricity production through electron transfer efficiency was noticed when acetate was used as a co-substrate. Apollon et al. (2023b) reported that using livestock urine as substrate resulted in a maximum PD and CD of 110.72±0.42 mW/m² and 277±0.04 mW/m², respectively.

Qi et al. (2018) suggested that sustainable energy could be generated by tapping on the potential of phototrophic bacteria (PTB). They further recorded an OCV and short–circuit current (ISC) of 0.96 V and 0.75 A/m² respectively, with a PTB. Several studies of pure or mixed culture of PTB and other electrogenic bacteria had been conducted: Geobacter metallireducens recorded a PD of 2.2 mW/m²; Rhodospirillum rubrum 1.25 W/m²; and Rhodopseudomonas palustris strain DX–1 2.7 W/m² (Xing et al., 2008).

Other MFC applications include energizing low–power consuming devices like small telemetry systems, wireless sensors, biosensors (for monitoring biological oxygen demand), and health condition monitors (Kumar et al., 2017). These developed devices include EcoBot I, EcoBot II, Slugbot, Gastrobot, and Chew Chew (Du et al., 2007; David and Higson, 2007).

Several modifications to enhance the performance of MFCs has been conducted; however, despite MFCs’ promising potential, the ideal low–power generation is a significant setback to its applicability (Wang et al., 2013). PTB uses light energy to degrade waste substances while generating electricity, but there is a paucity of research on harnessing and analyzing the electrochemical potential of PTB (Fischer, 2018). This study will assess and improve the electrochemical potential of a PTB by fabricating a 3D anode electrode using glucose as the substrate in a two–chamber MFC.

MATERIALS AND METHODS

A pure culture of the PTB previously isolated in another study was sub-cultured on agar and broth of 112 medium. Streak–plating method was adopted for the inoculation on an agar medium. The medium composition (L⁻¹) was 1.0 g of K₂HPO₄, 0.5 g of MgSO₄, 10.0 g of yeast extract, and 20.0 g of agar (only for agar medium). The culturing was done at 28–35℃ and 7.0–7.2 of temperature and pH, respectively, under an anaerobic, aseptic, and light mode for optimal growth. The bacterial growth (optical density) was monitored using a spectrophotometer at a wavelength of 660 nm (OD₆₆₀) (Kodama et al., 2012).

Morphological Study of the PTB

The gram staining technique and motility test were carried out according to the standard procedure to characterise the PTB in this research. According to Tripathi and Sapra
(2023), a single colony of the PTB was picked from a 24-hour agar plate using a sterile inoculating loop, smeared on a cleaned glass slide, and left to dry. The smear was first flooded with crystal violet stain for 60 secs, followed by gram’s iodine for 60 secs, a few drops of ethanol at 45⁰, and then safranin for 60 secs. Distilled water was used to wash the slide at each step before applying another reagent. After adding a drop of immersion oil and coverslip, a compound microscope was used to observe the slides.

The hanging drop method was applied to examine the bacterial motility. A single colony from the agar plate was picked using a sterile inoculating loop and placed in distilled water on a cleaned circular depression glass slide. Paraffin wax was used to surround the circular depression on the glass slide. With the help of a coverslip, the glass slide was fixed in an upside-down orientation on the compound microscope, and the bacterium was examined (Kodama et al., 2012).

Biochemical Tests

The two biochemical tests, catalase and oxidase, were conducted. One drop of PTB broth was set on a cleaned, labelled glass slide for the catalase test. While hydrogen peroxide (30% v/v) was the catalase reagent, a few drops were released on the bacteria and left for a few minutes to observe. Bubbles indicate a positive result, and their absence is negative (Azhary et al., 2020).

The oxidase test was done with a commercial oxidase reagent, Becton (5 mL). One drop of the bacterial suspension was also placed on a cleaned labelled glass slide. The reagent was deposited according to the manufacturer’s instructions on the bacterial drop and allowed for a few minutes before analysing. The result is positive when the color changes to dark purple within 10 secs, delayed positive when it takes 60-90 secs, and negative when it does not change longer than 120 secs (Dharmappa et al., 2022).

Polymerase Chain Reaction (PCR)

Colony PCR further studied the bacterium to amplify the 16S rRNA sequence. A single colony from an agar plate containing pure cultures of the PTB isolate was picked and suspended in 15 μL of distilled water, boiled in a PCR thermocycler for 10 mins, and centrifuged at 20,000 rpm for 5 mins. 1 μL of the supernatant was used as the DNA template.

The universal primers 27F (5′-AGAGTTTGTATCMTGGCTCAG-3′) and 1492R (5′-GGTTACCTTGTAGACTT-3′) were used in this study. The PCR mixture contained 50 μL per reaction: 25 μL of GoTaq green, 1 μL of each of 27F and 1492R primers, 1 μL of DNA template, and 22 μL of ultrapure water. The PCR amplification was done in the GeneAmp PCR System 9700 thermocycler (Perkin Elmer, GA9700) with an optimised amplification program of 1 cycle at 95°C for 2 mins, followed by 40 cycles of denaturation at 95°C for 45 secs, annealing at 55°C for 45 secs, extension at 72°C for 30 secs, and a final extension of 1 cycle at 72°C for 5 mins (Sharma et al., 2016).

Gel electrophoresis of the PCR product was performed using 1% gel prepared by weighing 1.5 g of agarose powder and mixing it with 250 mL of Tris–acetate buffer (TAE buffer). The solution was microwaved for 2–3 mins, cooled to room temperature, and 5 μL of ethidium bromide was added before pouring it into the comb tray to solidify. After solidification, 1,500 bp was deposited as the loading band in the first well. The electrophoresis machine was set for 30 mins at 80-120 V, and the products were visualised in the darkroom using ultraviolet (UV) light (Lee et al., 2012).

The PCR product was sequenced via the Sanger sequencing technique with initial ribonuclease treatment followed by DNA purification. The 16S rRNA gene sequence was obtained through Amplicon sequencing using NGS–7000 series. The obtained sequence was extracted using Sequence Scanner (SeqScan v2). The GenBank database blasted the extracted sequence using Nucleotide BLAST (blastN). Using the neighbour–joining tree method, a phylogenetic tree was constructed based on 100 related 16S rRNA gene sequences using molecular evolutionary genetics analysis (MEGA 11) (Tamura et al., 2021).

Preparation of Electrode

Graphene Oxide (GO) Preparation

The Hummer method was used to prepare the GO from graphite powder, as adopted by Wang et al. (2013). The graphite powder weighing 50 mg was mixed with 23 mL and 10 mL of concentrated H2SO4 and concentrated HNO3, respectively, in a cooled ice bath container, followed by a slow addition of 3 g KMnO4. The
solution was heated for 3 h at 35°C and then diluted with 40 mL of deionised water, followed by 200 mL of deionised water 12 h later, and subsequently, 3 mL of 30% v/v H₂O₂ was slowly added. The solution was centrifuged at 1,500 rpm for 30 mins, and the residue of graphite oxide was isolated and washed with deionized water before being resuspended in deionized water (500 mL). The aqueous solution was sonicated for 2 h to form layered GO sheets.

Reduced Graphene Oxide (rGO) Sheets Deposition on Nickel Foam (Ni)

Initially, nickel (Ni) foam was washed with diluted concentrated H₂SO₄ and placed in an oven at 60°C for 12 h. The Ni was then transferred into a 100 mL Teflon liner (hydrothermal reactor) filled with 50 mL of the sonicated aqueous GO solution, and the reactor was settled in an oven at 120°C for 5 h. After 5 h, it was allowed to cool down to room temperature, washed with deionized water, and air-dried. To increase the electrical conductivity and firmness of the rGO–Ni, annealing was performed at 400°C for 30 mins. The concentration of rGO in the solution was measured at 1 mg/mL (Wang et al., 2013).

Field Emission Scanning Electron Microscopy (FESEM)

The FESEM of Ni and rGO–Ni was conducted at the Department of Physics, Universiti Malaya, using the Joel (JSM–7600F) microscopy model. The secondary electron image (SEI) resolution and X-ray beam diameter were adjusted at 5 kV and 10 mm using the EDX–Oxford instrument. Images were obtained at different magnifications and diameters (Lawan et al., 2018).

Cyclic Voltammetry (CV)

The CVs were performed in a three-electrode set-up using an electrochemical workstation potentiostat: Eco Chemie (Autolab PGSTAT302). Ni and rGO–Ni were used as the working electrodes (WEs), Ag/AgCl as the reference electrode (RE), and the platinum wire as the counter electrode (CE). The WEs were cleaned with ethanol and washed with deionised water before use. The scan rate was set at 50 mV/s, and the potential ranges at 0–1 V. Measurements were taken repeatedly in 100 mM phosphate buffer solution (PBS) of 7.0 pH at room temperature and atmospheric pressure under aerobic and anaerobic conditions with and without substrate (glucose). Voltammograms were obtained via the computer software General Purpose Electrochemical System (GPES) v4.9 (Sharma et al., 2016).

Constructing the Microbial Fuel Cell (MFC)

A 100 mL two-chamber (50 mL each chamber) glassware MFC was used, as shown in Figure 2. Nafion NR–212 membrane was employed as the proton exchange membrane (PEM). The Nafion was pre-treated, as Rahmani et al. (2020) and Cheng et al. (2016) described. Originally, the membrane was boiled four times in distilled water, 30% v/v H₂O₂, 0.5M H₂SO₄, and finally, distilled water again at 80°C for 1 h each. The membrane was washed consecutively after every boiling before being put aside in distilled water for further use.

The electrodes, anodes (Ni/rGO–Ni), and cathode (carbon cloth) were 1×2 cm each. The anodes were subjected to pre-treatment to enhance electron transfer and microbial adhesion, as Cheng et al. (2016) mentioned. The electrodes were immersed in two chemicals: 30% v/v isopropanol overnight and 30% v/v H₂O₂ for 24 hours. The electrodes were washed with distilled water and dried thoroughly at 100°C at first and between the chemicals’ immersion. Contrary to the anode, as Rahmani et al. (2020) stated, the cathode was pre-treated by soaking in acetone for 20 mins and later boiling in 0.1 M of HCl for 15 mins.

The glucose–fed MFC was sterilised with ethanol first. Glucose (100 mM) was the substrate used. The anolyte consisted of the substrate, bacterial stock at OD₆₆₀ nm = 2, and 100 mM of PBS at 7.0 pH, while the catholyte was composed of 100 mM potassium ferricyanide. Although nitrogen gas was used to flush out oxygen in the anode chamber, so an anaerobic condition is maintained, the cathode chamber was left aerobic, as demonstrated in Figure 2. The MFC was connected to an external load of 1 kΩ for all experiments, the electrochemical station, and measurements were taken from GPES (v4.9) software. Experiments were run in batch mode, and each experiment was conducted twice to enhance data accuracy. The power was calculated using P=I*V, and the power densities were calculated based on the anode area (2 cm² or 0.0002 m²) (Haruna et al., 2023).
RESULTS

Morphological Study

The bacterial colonies on the agar 24 h later formed a smooth, circular, milky color, less than 1 mm in diameter. On the contrary, the bacterial broth showed a dense reddish color at day 10, while the OD measured at 660 nm was 2.

The observed microscopy of the gram staining reaction showed it to be a chain-like red bacterium with an approximate size of 1.2×1.4 um. The motility examination of the bacterium revealed no movement on the slide completely. In addition, there was no flagellum or sporulating seen.

Biochemical Tests

Consequently, both the slides of catalase and oxidase tests, when observed, produced no reaction after 2 mins. The absence of air bubbles on the catalase slide showed the bacterium does not produce catalase enzyme hence it is catalase-negative. This further indicated that the bacterium does not undergo aerobic respiration, confirming it to be anaerobic. It was also observed that the presence of oxygen affects the growth rate of the bacterium, ultimately influencing the performance of the MFC.

Polymerase Chain Reaction (PCR)

The gel electrophoresis results showed an estimated 1,200 bp band of the PCR product under UV light. A partial 16S rRNA sequence of 1,211 bp was acquired from the extracted sequence. The blast N analysis of the 16S rRNA partial sequence in the GenBank database revealed 100 significant strains of microorganisms with similarities ranging from 84.10% to 98.76%.

Of the 100 strains of 16S rRNA partial sequences deposited in GenBank, 13 have the highest similarities (>90%), all from the genus Dysgonomonas. Table 1 shows that Dysgonomonas oryzarvi Dy73 has the closest (98.76%) sequence similarity, and Dysgonomonas capnocytophagoides CDC F9047 (90.36%) has the farthest similarity. This analysis suggested that the bacterium belongs to the genus Dysgonomonas.
Table 1: The 13 strains with the highest similarities (>90%) to the PTB

<table>
<thead>
<tr>
<th>S/No.</th>
<th>Strains</th>
<th>Homology (%)</th>
</tr>
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<tbody>
<tr>
<td>1.</td>
<td><em>Dysgonomonas oryzarvi</em> Dy73</td>
<td>98.76</td>
</tr>
<tr>
<td>2.</td>
<td><em>Dysgonomonas mossii</em> DSM 22836 JCM 16699</td>
<td>98.60</td>
</tr>
<tr>
<td>3.</td>
<td><em>Dysgonomonas mossii</em> CCUG 43457</td>
<td>98.51</td>
</tr>
<tr>
<td>4.</td>
<td><em>Dysgonomonas termitidis</em> JCM 30204</td>
<td>93.96</td>
</tr>
<tr>
<td>5.</td>
<td><em>Dysgonomonas gadei</em> ATCC BAA–286 JCM 16698</td>
<td>93.64</td>
</tr>
<tr>
<td>7.</td>
<td><em>Dysgonomonas massiliensis</em> Marseille–P4356</td>
<td>93.17</td>
</tr>
<tr>
<td>8.</td>
<td><em>Dysgonomonas hofsadii</em> JCM 17038</td>
<td>92.08</td>
</tr>
<tr>
<td>9.</td>
<td><em>Dysgonomonas hofsadii</em> MX 1040</td>
<td>92.00</td>
</tr>
<tr>
<td>10.</td>
<td><em>Dysgonomonas alginitilvica</em> HUA–2</td>
<td>91.23</td>
</tr>
<tr>
<td>11.</td>
<td><em>Dysgonomonas capnocytophagoides</em> JCM 16697</td>
<td>91.10</td>
</tr>
<tr>
<td>12.</td>
<td><em>Dysgonomonas macrotermitis</em> Dys–CH1</td>
<td>90.90</td>
</tr>
<tr>
<td>13.</td>
<td><em>Dysgonomonas capnocytophagoides</em> CDC F9047</td>
<td>90.36</td>
</tr>
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Figure 3. Phylogenetic tree with the PTB’s inquiry ID as 4599196 CTRL A.

The bacterium's phylogenetic tree (Figure 3) was created in the molecular evolutionary genetics analysis (MEGA 11) using the neighbour-joining method as mentioned in the methods. The PTB has not been reported in MFCs before, and this could be due to the absence of cytochrome enzyme, which is responsible for electron transfer during bacterial metabolism (Robb, 2020).

Electrode Preparation

The change in Ni color to black after rGO deposition indicated the effectiveness of the process (Figure 4a and b). Despite the annealing at 400°C for 5 h, the flexibility remained intact with no fragmentation when bent (Figure 4c). This makes the electrode apt for any MFC type and scaling-up application.

As revealed by the SEM images, the Ni was completely embedded with the rGO (Figure 5e and f), while Figure 5d confirmed multi-layered GO sheets in a continuous 3D scaffold with pores. These pores accommodate more bacterial adhesion, which increases electrochemical
activities, thereby enhancing the overall performance of the MFC.

As shown in Figure 4b, the change in the colour of Ni demonstrated rGO deposition. Further examination through SEM images proved the complete embedment of rGO onto Ni to form the 3D rGO–Ni electrode.

The CV analyses under the anaerobic and aerobic conditions with or without glucose showed oxidation and reduction peaks against the saturated RE: Ag/AgCl. The aerobic CVs with and without substrate revealed uncleared curves for Ni and rGO–Ni. In addition, the least maximum redox peaks were obtained in the aerobic treatment without substrate: 0.07 nA(Ni) and 33.6 nA(rGO–Ni) (Figure 6a and e). Cleared curves for anaerobic with and without substrate were seen.

Figure 4. (a) Ni before rGO deposition, (b) Ni after deposition and annealing of rGO to form the 3D rGO–Ni electrode, and (c) bending of the rGO–Ni, which confirmed the intactness of its flexibility without fragmentation.

Figure 5. SEM images of the Ni before deposition and annealing (a–c) and after deposition and annealing (d–f).
Figure 6. CV measurements of Ni (a-d) and rGO–Ni (e-f) were measured. Both measurements of electrodes were taken under aerobic conditions without substrate (a and e), aerobic with substrate (b and f), anaerobic without substrate (c and g), and anaerobic with substrate (d and h).

Figure 7. The short-circuit current generated over 11 days in a batch mode MFC.
A maximum OCV of 0.99 V was measured from rGO–Ni while Ni recorded 0.34 V (Figure 8). Although this is four times lower than that obtained from the microfluid–MFC configuration by Shirkosh et al. (2022); however, it is also 100-fold higher compared to the 0.037 mV reported by Tou et al. (2019). This could result from the potential of the electrogenic bacterium and the enhanced 3D anode. Similarly, the rGO–Ni and Ni electrodes also exhibited a maximum ISC of 0.99 mA and 0.25 mA, respectively (Figure 8).

**Figure 8.** The open circuit voltage is measured in a batch mode MFC.

**DISCUSSION**

The absence of retaining stain made it gram–negative, as discussed by several researchers. Neither a flagellum nor sporulation was seen. It has a chain–like coccus with circular colonies of less than 1 mm. This aligned with the characteristics of a non–motile, gram–negative bacterium, as Moyes et al. (2009) indicated.

The catalase result showed the bacterium to be anaerobic. Further observations revealed the presence of oxygen to affect the bacterial growth slightly. In the case of oxidase result, the bacterium cannot produce cytochrome oxidase enzyme, which enhances the electrochemical activity of electrogenic bacteria (Sharma et al., 2016). This also indicates the presence or absence of the enzyme influences the power generation as stated by Sonaware et al. (2017). The molecular technique attested that the bacterium was likely *Dysgonomonas oryzarvi* Dy73, with 98.76% similarities as the morphological features, biochemical tests, and molecular study indicated (Kodama et al., 2012).

The CVs conducted on Ni (Figure 6a–d) and rGO–Ni (Figure 6e–f) confirmed an enhanced electrochemical potential of the rGO–Ni electrode compared to the Ni. Not only did the CVs show that the bacterium is electrochemically active, but they also proved the effect of aerobic conditions on the electrochemical activity of the bacterium. Although small redox peaks were recorded in aerobic conditions without any substrate (Figure 6a and e), no changes were seen for the CV curves of both electrodes in anaerobic conditions with substrate (Figure 6d and h) except that the curves were clearer and attained the maximum peak in Figure 5h. As seen in Figure 5, the maximum peaks shared a similar pattern and were recorded from anaerobic treatments with a substrate, rGO–Ni (8.88 mA), followed by anaerobic with no substrate, rGO–Ni.
(6.80 mA), anaerobic with a substrate, specifically Ni (5.95 mA), and anaerobic without substrate, also in the Ni treatment (4.54 mA). In summary, the fabricated 3D electrode accommodates more bacterial adhesion, as Wang et al. (2013) suggested, thereby increasing the power density produced by the MFC. Additionally, the material used, Ni foam, plays a significant role in being unharmed material and enhancing the bacterium's electrochemical productivity.

The maximum PD values of approximately 4.9 W/m² and 0.58 W/m² were calculated for rGO–Ni and Ni electrodes. This is significantly higher than studies conducted by Eslami et al. (2023) and Tou et al. (2019) that generated a PD of 0.7 mW/m² when graphite rod was used with Saccharomyces cerevisiae and 744 uW/m² using Chlophytum comosum, respectively. However, compared to the 3D GN–Co₃O₄ used by Tan et al. (2018), a notable maximum PD of 578±10 mW/m² was recorded. This resulted from the modification and electrode material used, as Chen et al. (2014) and Hou et al. (2014) pointed out.

Approximately one-and-a-half folds of OCV, four folds of ISC, and six folds of PD were obtained from rGO–Ni compared to the Ni electrode, proving the effectiveness and efficiency of the continuous 3D scaffold electrode in increasing microbial colonisation as well as enhancing the electrochemical performance of the overall MFC. Furthermore, the cell ran for 11 days before the current started to decline progressively, suggesting the depletion of nutrients and possibly the removal of the organic matter in the anolyte. It was also deciphered that the cell could run for more than two weeks while electrochemically functioning.

CONCLUSION

The phototrophic bacterium in this study was a gram-negative, non-motile coccus in chains that is a facultatively anaerobic bacterium that belongs to the genus Dysgonomonas. Based on the CV, OCV, ISC, and PD values of the two electrodes, Ni (0.58 W/m²) and rGO–Ni (4.9 W/m²), this study established that the fabrication of a 3D rGO–Ni electrode increases microbial adhesion, colonisation, as well as the overall power density generated by the D. oryzae-rvi Dy73in the MFC providing more applicable, economically viable, and sustainable alternative to bioelectricity generation. More importantly, this study demonstrated that 3D electrodes can further amplify MFC performance and could be scaled industrially. Further analyses to understand this bacterium's exact mechanism of exocellular electron transfer are recommended for future study.

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